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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/023,523

DATE: 02/11/2002
 TIME: 08:34:36

Input Set : N:\Crf3\RULE60\10023523.raw
 Output Set: N:\CRF3\02112002\J023523.raw

1 <110> APPLICANT: Lees, Ann M.
 2 Lees, Robert S.
 3 Law, Simon W.
 4 Arjona, Anibal A.
 5 <120> TITLE OF INVENTION: NOVEL LOW DENSITY LIPOPROTEIN BINDING
 6 PROTEINS AND THEIR USE IN DIAGNOSING AND TREATING
 7 ATHEROSCLEROSIS
 8 <130> FILE REFERENCE: 10797-004001
 9 <140> CURRENT APPLICATION NUMBER: US/10/023,523
 10 <141> CURRENT FILING DATE: 2001-12-17
 11 <150> PRIOR APPLICATION NUMBER: US/09/616,289
 12 <151> PRIOR FILING DATE: 2000-07-14
 13 <150> PRIOR APPLICATION NUMBER: US 09/517,849
 14 <151> PRIOR FILING DATE: 2000-03-02
 15 <150> PRIOR APPLICATION NUMBER: US 08/979,608
 16 <151> PRIOR FILING DATE: 1997-11-26
 17 <150> PRIOR APPLICATION NUMBER: US 60/031,930
 18 <151> PRIOR FILING DATE: 1996-11-27
 19 <150> PRIOR APPLICATION NUMBER: US 60/048,547
 20 <151> PRIOR FILING DATE: 1997-06-03
 21 <160> NUMBER OF SEQ ID NOS: 53
 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 151
 26 <212> TYPE: PRT
 27 <213> ORGANISM: Oryctolagus cuniculus
 28 <400> SEQUENCE: 1
 29 Met Ser Lys Asn Thr Val Ser Ser Ala Arg Phe Arg Lys Val Asp Val
 30 1 5 10 15
 31 Asp Glu Tyr Asp Glu Asn Lys Phe Val Asp Glu Glu Asp Gly Gly Asp
 32 20 25 30
 33 Gly Gln Ala Gly Pro Asp Glu Gly Glu Val Asp Ser Cys Leu Arg Gln
 34 35 40 45
 35 Gly Asn Met Thr Ala Ala Leu Gln Ala Ala Leu Lys Asn Pro Pro Ile
 36 50 55 60
 37 Asn Thr Arg Ser Gln Ala Val Lys Asp Arg Ala Gly Ser Ile Val Leu
 38 65 70 75 80
 39 Lys Val Leu Ile Ser Phe Lys Ala Gly Asp Ile Glu Lys Ala Val Gln
 40 85 90 95
 41 Ser Leu Asp Arg Asn Gly Val Asp Leu Leu Met Lys Tyr Ile Tyr Lys
 42 100 105 110
 43 Gly Phe Glu Ser Pro Ser Asp Asn Ser Ser Ala Val Leu Leu Gln Trp
 44 115 120 125

ENTERED

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45 His Glu Lys Ala Leu Ala Ala Gly Gly Val Gly Ser Ile Val Arg Val
46 130 135 140
47 Leu Thr Ala Arg Lys Thr Val
48 145 150
50 <210> SEQ ID NO: 2
51 <211> LENGTH: 317
52 <212> TYPE: PRT
53 <213> ORGANISM: Oryctolagus cuniculus
54 <220> FEATURE:
55 <221> NAME/KEY: VARIANT
56 <222> LOCATION: (1)...(317)
57 <223> OTHER INFORMATION: Xaa = Any Amino Acid
58 <400> SEQUENCE: 2
W--> 59 Asp Cys Arg Ser Ser Asn Asn Arg Xaa Pro Lys Gly Gly Ala Ala
60 1 5 10 15
61 Arg Ala Gly Gly Pro Ala Arg Pro Val Ser Leu Arg Glu Val Val Arg
62 20 25 30
63 Tyr Leu Gly Gly Ser Ser Gly Ala Gly Gly Arg Leu Thr Arg Gly Arg
64 35 40 45
65 Val Gln Gly Leu Leu Glu Glu Ala Ala Ala Arg Gly Arg Leu Glu
66 50 55 60
67 Arg Thr Arg Leu Gly Ala Leu Ala Leu Pro Arg Gly Asp Arg Pro Gly
68 65 70 75 80
69 Arg Ala Pro Pro Ala Ala Ser Ala Arg Ala Ala Arg Asn Lys Arg Ala
70 85 90 95
71 Gly Glu Glu Arg Val Leu Glu Lys Glu Glu Glu Glu Glu Glu Glu
72 100 105 110
73 Asp Asp Glu Asp Asp Asp Asp Val Val Ser Glu Gly Ser Glu Val
74 115 120 125
75 Pro Glu Ser Asp Arg Pro Ala Gly Ala Gln His His Gln Leu Asn Gly
76 130 135 140
77 Gly Glu Arg Gly Pro Gln Thr Ala Lys Glu Arg Ala Lys Glu Trp Ser
78 145 150 155 160
79 Leu Cys Gly Pro His Pro Gly Gln Glu Glu Gly Arg Gly Pro Ala Ala
80 165 170 175
81 Gly Ser Gly Thr Arg Gln Val Phe Ser Met Ala Ala Leu Ser Lys Glu
82 180 185 190
83 Gly Gly Ser Ala Ser Ser Thr Thr Gly Pro Asp Ser Pro Ser Pro Val
84 195 200 205
85 Pro Leu Pro Pro Gly Lys Pro Ala Leu Pro Gly Ala Asp Gly Thr Pro
86 210 215 220
87 Phe Gly Cys Pro Ala Gly Arg Lys Glu Lys Pro Ala Asp Pro Val Glu
88 225 230 235 240
89 Trp Thr Val Met Asp Val Val Glu Tyr Phe Thr Glu Ala Gly Phe Pro
90 245 250 255
91 Glu Gln Ala Thr Ala Phe Gln Glu Gln Glu Ile Asp Gly Lys Ser Leu
92 260 265 270
93 Leu Leu Met Gln Arg Thr Asp Val Leu Thr Gly Leu Ser Ile Arg Leu
94 275 280 285

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95      Gly Pro Ala Leu Lys Ile Tyr Glu His His Ile Lys Val Leu Gln Gln
96          290           295           300
97      Gly His Phe Glu Asp Asp Asp Pro Glu Gly Phe Leu Gly
98          305           310           315
100 <210> SEQ ID NO: 3
101 <211> LENGTH: 232
102 <212> TYPE: PRT
103 <213> ORGANISM: Oryctolagus cuniculus
104 <400> SEQUENCE: 3
105      Ala Ser Ala Arg Ala Ala Arg Asn Lys Arg Ala Gly Glu Glu Arg Val
106          1             5             10            15
107      Leu Glu Lys Glu Glu Glu Glu Glu Glu Asp Asp Glu Asp Asp
108          20            25            30
109      Asp Asp Asp Val Val Ser Glu Gly Ser Glu Val Pro Glu Ser Asp Arg
110          35            40            45
111      Pro Ala Gly Ala Gln His His Gln Leu Asn Gly Gly Glu Arg Gly Pro
112          50            55            60
113      Gln Thr Ala Lys Glu Arg Ala Lys Glu Trp Ser Leu Cys Gly Pro His
114          65            70            75            80
115      Pro Gly Gln Glu Glu Gly Arg Gly Pro Ala Ala Gly Ser Gly Thr Arg
116          85            90            95
117      Gln Val Phe Ser Met Ala Ala Leu Ser Lys Glu Gly Gly Ser Ala Ser
118          100           105           110
119      Ser Thr Thr Gly Pro Asp Ser Pro Ser Pro Val Pro Leu Pro Pro Gly
120          115           120           125
121      Lys Pro Ala Leu Pro Gly Ala Asp Gly Thr Pro Phe Gly Cys Pro Ala
122          130           135           140
123      Gly Arg Lys Glu Lys Pro Ala Asp Pro Val Glu Trp Thr Val Met Asp
124          145           150           155           160
125      Val Val Glu Tyr Phe Thr Glu Ala Gly Phe Pro Glu Gln Ala Thr Ala
126          165           170           175
127      Phe Gln Glu Gln Glu Ile Asp Gly Lys Ser Leu Leu Leu Met Gln Arg
128          180           185           190
129      Thr Asp Val Leu Thr Gly Leu Ser Ile Arg Leu Gly Pro Ala Leu Lys
130          195           200           205
131      Ile Tyr Glu His His Ile Lys Val Leu Gln Gln Gly His Phe Glu Asp
132          210           215           220
133      Asp Asp Pro Glu Gly Phe Leu Gly
134          225           230
136 <210> SEQ ID NO: 4
137 <211> LENGTH: 252
138 <212> TYPE: PRT
139 <213> ORGANISM: Oryctolagus cuniculus
140 <400> SEQUENCE: 4
141      Thr Arg Leu Gly Ala Leu Ala Leu Pro Arg Gly Asp Arg Pro Gly Arg
142          1             5             10            15
143      Ala Pro Pro Ala Ala Ser Ala Arg Ala Ala Arg Asn Lys Arg Ala Gly
144          20            25            30
145      Glu Glu Arg Val Leu Glu Lys Glu Glu Glu Glu Glu Asp

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```

146      35          40          45
147 Asp Glu Asp Asp Asp Asp Val Val Ser Glu Gly Ser Glu Val Pro
148      50          55          60
149 Glu Ser Asp Arg Pro Ala Gly Ala Gln His His Gln Leu Asn Gly Gly
150      65          70          75          80
151 Glu Arg Gly Pro Gln Thr Ala Lys Glu Arg Ala Lys Glu Trp Ser Leu
152      85          90          95
153 Cys Gly Pro His Pro Gly Gln Glu Glu Gly Arg Gly Pro Ala Ala Gly
154      100         105         110
155 Ser Gly Thr Arg Gln Val Phe Ser Met Ala Ala Leu Ser Lys Glu Gly
156      115         120         125
157 Gly Ser Ala Ser Ser Thr Thr Gly Pro Asp Ser Pro Ser Pro Val Pro
158      130         135         140
159 Leu Pro Pro Gly Lys Pro Ala Leu Pro Gly Ala Asp Gly Thr Pro Phe
160      145         150         155         160
161 Gly Cys Pro Ala Gly Arg Lys Glu Lys Pro Ala Asp Pro Val Glu Trp
162      165         170         175
163 Thr Val Met Asp Val Val Glu Tyr Phe Thr Glu Ala Gly Phe Pro Glu
164      180         185         190
165 Gln Ala Thr Ala Phe Gln Glu Glu Ile Asp Gly Lys Ser Leu Leu
166      195         200         205
167 Leu Met Gln Arg Thr Asp Val Leu Thr Gly Leu Ser Ile Arg Leu Gly
168      210         215         220
169 Pro Ala Leu Lys Ile Tyr Glu His His Ile Lys Val Leu Gln Gln Gly
170      225         230         235         240
171 His Phe Glu Asp Asp Asp Pro Glu Gly Phe Leu Gly
172      245         250
174 <210> SEQ ID NO: 5
175 <211> LENGTH: 557
176 <212> TYPE: PRT
177 <213> ORGANISM: Oryctolagus cuniculus
178 <400> SEQUENCE: 5
179 Met Lys Asn Gln Asp Lys Lys Asn Gly Ala Ala Lys Gln Pro Asn Pro
180      1           5           10          15
181 Lys Ser Ser Pro Gly Gln Pro Glu Ala Gly Ala Glu Gly Ala Gln Gly
182      20          25          30
183 Arg Pro Gly Arg Pro Ala Pro Ala Arg Glu Ala Glu Gly Ala Ser Ser
184      35          40          45
185 Gln Ala Pro Gly Arg Pro Glu Gly Ala Gln Ala Lys Thr Ala Gln Pro
186      50          55          60
187 Gly Ala Leu Cys Asp Val Ser Glu Glu Leu Ser Arg Gln Leu Glu Asp
188      65          70          75          80
189 Ile Leu Ser Thr Tyr Cys Val Asp Asn Asn Gln Gly Ala Pro Gly Glu
190      85          90          95
191 Asp Gly Val Gln Gly Glu Pro Pro Glu Pro Glu Asp Ala Glu Lys Ser
192      100         105         110
193 Arg Ala Tyr Val Ala Arg Asn Gly Glu Pro Glu Pro Gly Thr Pro Val
194      115         120         125
195 Val Asn Gly Glu Lys Glu Thr Ser Lys Ala Glu Pro Gly Thr Glu Glu

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196	130	135	140
197	Ile Arg Thr Ser Asp Glu Val Gly Asp Arg Asp His Arg Arg Pro Gln		
198	145	150	155
199	Glu Lys Lys Lys Ala Lys Gly Leu Gly Lys Glu Ile Thr Leu Leu Met		160
200	165	170	175
201	Gln Thr Leu Asn Thr Leu Ser Thr Pro Glu Glu Lys Leu Ala Ala Leu		
202	180	185	190
203	Cys Lys Lys Tyr Ala Glu Leu Leu Glu Glu His Arg Asn Ser Gln Lys		
204	195	200	205
205	Gln Met Lys Leu Leu Gln Lys Lys Gln Ser Gln Leu Val Gln Glu Lys		
206	210	215	220
207	Asp His Leu Arg Gly Glu His Ser Lys Ala Ile Leu Ala Arg Ser Lys		
208	225	230	235
209	Leu Glu Ser Leu Cys Arg Glu Leu Gln Arg His Asn Arg Ser Leu Lys		240
210	245	250	255
211	Glu Glu Gly Val Gln Arg Ala Arg Glu Glu Glu Lys Arg Lys Glu		
212	260	265	270
213	Val Thr Ser His Phe Gln Met Thr Leu Asn Asp Ile Gln Leu Gln Met		
214	275	280	285
215	Glu Gln His Asn Glu Arg Asn Ser Lys Leu Arg Gln Glu Asn Met Glu		
216	290	295	300
217	Leu Ala Glu Arg Leu Lys Lys Leu Ile Glu Gln Tyr Glu Leu Arg Glu		
218	305	310	315
219	Glu His Ile Asp Lys Val Phe Lys His Lys Asp Leu Gln Gln Leu		320
220	325	330	335
221	Val Asp Ala Lys Leu Gln Gln Ala Gln Glu Met Leu Lys Glu Ala Glu		
222	340	345	350
223	Glu Arg His Gln Arg Glu Lys Asp Phe Leu Leu Lys Glu Ala Val Glu		
224	355	360	365
225	Ser Gln Arg Met Cys Glu Leu Met Lys Gln Gln Glu Thr His Leu Lys		
226	370	375	380
227	Gln Gln Leu Ala Leu Tyr Thr Glu Lys Phe Glu Glu Phe Gln Asn Thr		
228	385	390	395
229	Leu Ser Lys Ser Ser Glu Val Phe Thr Thr Phe Lys Gln Glu Met Glu		400
230	405	410	415
231	Lys Met Thr Lys Ile Lys Lys Leu Glu Lys Glu Thr Thr Met Tyr		
232	420	425	430
233	Arg Ser Arg Trp Glu Ser Ser Asn Lys Ala Leu Leu Glu Met Ala Glu		
234	435	440	445
235	Glu Lys Thr Leu Arg Asp Lys Glu Leu Glu Gly Leu Gln Val Lys Ile		
236	450	455	460
237	Gln Arg Leu Glu Lys Leu Cys Arg Ala Leu Gln Thr Glu Arg Asn Asp		
238	465	470	475
239	Leu Asn Lys Arg Val Gln Asp Leu Ser Ala Gly Gly Gln Gly Pro Val		480
240	485	490	495
241	Ser Asp Ser Gly Pro Glu Arg Arg Pro Glu Pro Ala Thr Thr Ser Lys		
242	500	505	510
243	Glu Gln Gly Val Glu Gly Pro Gly Ala Gln Val Pro Asn Ser Pro Arg		
244	515	520	525

VERIFICATION SUMMARY

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TIME: 08:34:37

Input Set : N:\Crf3\RULE60\10023523.raw
Output Set: N:\CRF3\02112002\J023523.raw

L:59 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2

L:2730 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53